



## Multivariate ARIMA and ARIMA-X Analysis

Package 'marima'

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# Package ‘marima’

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arma.filter	<i>arma.filter</i>
-------------	--------------------

---

## Description

Filtering of (kvar-variate) time series with marima type model.

Calculation of residuals and filtered values of timeseries using a marima model.

## Usage

```
arma.filter(series = NULL, ar.poly = array(diag(kvar), dim = c(kvar, kvar, 1)), ma.poly = array(diag(kvar), dim = c(kvar, kvar, 1)), means = 1)
```

## Arguments

series	matrix holding the kvar by n multivariate timeseries (if (kvar > n) the series is transposed and a warning is given).
ar.poly	(kvar,kvar,p+1) array containing autoregressive matrix polynomial model part. If the filtering is to be performed for undifferenced data when the analysis (in marima) was done for differenced data, the input array ar.poly should incorporate the ar-representation of the differencing operation (using, for example: <code>ar.poly &lt;- pol.mul(ar.estimate, dif.poly, L = ( dim(ar.estimate)[3]+dim(dif.poly)[3]))</code> , where 'dif.poly' was obtained when differencing the time series (using <code>define.dif</code> ) before analysing it with marima (giving the <code>ar.estimate</code> ) .
ma.poly	(kvar,kvar,q+1) array containing moving average matrix polynomial model part. If a leading unity matrix is not included in the ar- and/or the ma-part of the model this is automatically taken care of in the function (in that case the dimensions of the model arrays used in <code>arma.filter()</code> are, respectively, (kvar,kvar,p+1) and (kvar,kvar,q+1)).
means	vector (length = kvar) indicating whether means are subtracted or not (0/1). Default : means=1 saying that all means are subtracted (equivalent to means = c(1,1,...,1)).

## Value

estimates = estimated values for input series

residuals = corresponding residuals

averages = averages of variables in input series

mean.pattern = pattern of means as used in filtering

## Examples

```
library(marima)
data(austr)
series<-t(austr)[,1:90]
# Define marima model
Model5 <- define.model(kvar=7,ar=1,ma=1,rem.var=1,reg.var=6:7)

# Estimate marima model
Marima5 <- marima(series,Model5$ar.pattern,Model5$ma.pattern,penalty=1)

# Calculate residuals by filtering
Resid <- arma.filter(series,Marima5$ar.estimates,
  Marima5$ma.estimates)
# Compare residuals

plot(Marima5$residuals[2,4:89], Resid$residuals[2,5:90],
  xlab='marima residuals', ylab='arma.filter residuals')
```

---

arma.forecast

*arma.forecast*

---

## Description

Forecasting of (multivariate) time series of marima type using marima type model.

## Usage

```
arma.forecast(series = NULL, marima = NULL, nstart = NULL, nstep = 1,
  dif.poly = NULL)
```

## Arguments

**series** = matrix holding the kvar-variate timeseries. The series is assumed to have the same format as the timeseries analysed by marima BEFORE differencing (if differencing was used via define.dif) (the length, though, does not need to be the same but can be shorter or longer). Results from estimating the model (for the differenced data, if used) are assumed to be saved in the input-object 'marima' (see 'usage') by marima.

The series is assumed to have the total length=(nstart+nstep) (but it may be longer. In any case the forecasting is starting from nstart continuing to nstart+nstep. Future values already present or initialised, for example, as NAs are overwritten with the forecasted values.)

An example of a series prepared for forecasting is in the marima library: 'data(austr)': (see below, the example).

If future (independent) x-values for the forecasting are to be used these values must be supplied in 'series' at the proper places before calling 'arma.forecast(...)' (that is except the x-value(s) corresponding to the last prediction).

**marima** = the object holding the marima results to be used for the forecasting, that is an output object created by marima.

If the ar- and/or the ma-model do not include a leading unity matrix this is automatically taken care of in the function (in that case the dimensions of the model arrays used will be, respectively, (kvar,kvar,p+1) and (kvar,kvar,q+1)) after inserting the leading unity matrix (if the object 'marima' was produced by marima, this will automatically be OK).

**nstart** = starting point for forecasting (1st forecast values will be for time point t = nstart+1).

**nstep** = length of forecast (forecasts will be for time points nstart+1,...,nstart+nstep).

**dif.poly** (most often) output from the function define.dif holding the ar-representation of the differencing polynomial (define.dif\$dif.poly). If a differenced timeseries was analysed by marima the forecast-variance/covariance matrices are calculated for the aggregated (original) timeseries if 'dif.poly' is specified. If not, the forecast-variance/covariance matrices are calculated for the differenced time series. If forecasting is wanted for the original (not differenced) time series the 'dif.poly' created by define.dif must be specified.

### Value

forecasts = forecasted values following the nstart first values of the input series (at time points 'nstart+1,...,nstart+nstep'). The forecasted values will be (over-) written in the input series at the proper future positions (if relevant).

residuals = corresponding residuals for input series followed by nstep future residuals (all=0).

prediction.variances = (kvar,kvar,nstep) array containing prediction covariance matrices corresponding to the nstep forecasts.

nstart = starting point for prediction (1st prediction at point nstart+1).

nstep = length of forecast

### Examples

```
library(marima)
data(austr)
series<-t(austr)
Model5 <- define.model(kvar=7, ar=1, ma=1, rem.var=1, reg.var=6:7)
Marima5 <- marima(series[, 1:90], Model5$ar.pattern, Model5$ma.pattern,
  penalty=1)
nstart <- 90
nstep <- 10
Forecasts <- arma.forecast(series=series, marima=Marima5,
  nstart=nstart, nstep=nstep)
Year<-series[1, 91:100];
Predict<-Forecasts$forecasts[2, 91:100]
stdv<-sqrt(Forecasts$pred.var[2, 2, ])
upper.lim=Predict+stdv*1.645
lower.lim=Predict-stdv*1.645
Out<-rbind(Year, Predict, upper.lim, lower.lim)
print(Out)
```

```
# plot results:
plot(series[1, 1:100], Forecasts$forecasts[2, ], type='l', xlab='Year',
ylab='Rate of armed suicides', main='Prediction of suicides by firearms',
ylim=c(0.0, 4.1))
lines(series[1, 1:90], series[2, 1:90], type='p')
grid(lty=2, lwd=1, col='black')
Years<-2005:2014
lines(Years, Predict, type='l')
lines(Years, upper.lim, type='l')
lines(Years, lower.lim, type='l')
lines(c(2004.5, 2004.5), c(0.0, 2.0), lty = 2)
```

---

austr

*Data set for testing marima package (australian killings)*


---

## Description

Data set for testing marima package (australian killings)

## Usage

```
data(austr)
```

## Format

A data frame (austr) with 7 columns and 100 rows.

**price** Year

**suic.fire** Rate of suicides by firearms

**homi.fire** Rate of homicides by firearms

**suic.other** Rate of suicides by non firearms

**homi.other** Rate of homicides by non firearms

**leg** Legislation against firearms in effect

**acc.elg** Accumulated effect of legislation in years

---

 check.one

*check.one*


---

### Description

Function to check and insert leading unity matrix if NOT present.

### Usage

```
check.one(polyn = NULL)
```

### Arguments

polyn (k,k,...) matrix polynomium with or without leading unity matrix.

### Value

polyn (array) with a leading unity matrix being inserted if not present.

### Examples

```
set.seed(4711)
X<-array(rnorm(32),dim=c(4,4,2))
X<-check.one(X)
short.form(X)
```

---

 define.dif

*define.dif*


---

### Description

Function to generate and apply a differencing matrix polynomial (autoregressive form) defined by a pattern.

To be used before calling marima in order to difference the timeseries before the marima analysis. The averages of the variables in the time series are subtracted from the input series before differencing.

### Usage

```
define.dif(series = series, difference = NULL)
```

### Arguments

series = kvar-variate timeseries (kvar by n matrix).

difference = 2 by L matrix defining L differencing operations.

**Value**

y.dif = the differenced timeseries (the complete part)  
 y.lost = the first observations lost because of differencing  
 dif.poly = differencing polynomial array = c(kvar, kvar, ...) holding the autoregressive representation of the specified differencing  
 averages = the averages of the original series as they were subtracted before differencing  
 dif.series = the differenced series (y.lost followed by y.dif)

**Examples**

```
# Generate Y=series with 4 variables for illustration:
set.seed(4711)
Y<-matrix(round(100*rnorm(40)+10), nrow=4)

# Example 1: use of difference parameter: If
difference=c(2, 1, 2, 1, 3, 12)
difference
# the variable 2 is differenced
# twice, and variable 3 is differenced once with lag=12.

# Example 2:
poly <- define.dif(series=Y, difference=c(2, 1, 3, 1, 3, 1))
poly
# Generates a (4-variate) polynomial differencing array (with a leading
# unity matrix corresponding to lag=0, and (in the example) differencing
# of variable 2 for lag 1 and variable 3 for lag 1 but twice. Afterwards
# the series Y is differenced accordingly. Results in poly$series and
# poly$dif.poly .

# Example 3: Generation and application of multivariate differencing
# polynomial. Re-use the 4-variate time series and use the
# differencing polynomial (ar-form):
# var=1, dif=1, var=2, dif=6, and var=3 and 4, no differencing.
dif.y <-define.dif(Y, c(1, 1, 2, 6, 3, 0, 4, 0))
# Now dif.y contains the differenced series and the differencing
# polynomial. Print the generated polynomial in short form:
short.form(dif.y$dif.poly)
# Specifying no differencing (3, 0 and 4, 0) may be omitted:
dif.y <-define.dif(Y, c(1, 1, 2, 6))
dif.y

# Example 4:
y<-matrix(round(rnorm(1200)*100+50), nrow=6)
library(marima)
difference<-c(3, 2, 4, 0, 5, 0, 6, 7)
matrix(difference, nrow=2)
Y<-define.dif(y, difference=difference)
round(rowMeans(Y$dif.series), 2)
round(Y$averages, 2)
```



define.model

*define.model***Description**

Function to define multivariate arma model (indicator form) for marima.

**Usage**

```
define.model(kvar = 1, ar = 0, ma = 0, rem.var = 0, reg.var = 0,
             no.dep = NULL, print = 0, ar.fill = NULL, ar.rem = NULL,
             ma.fill = NULL, ma.rem = NULL, indep = NULL)
```

**Arguments**

kvar	= dimension of time series
ar	= autoregresssion definition. For example ar=c(1, 2, 12) will generate autoregression at lags 1, 2 and 12.
ma	= moving average definition. Works like ar. If ma=c(1, 2) moving average terms at lags 1 and 2 are defined.
rem.var	= no. of variable(s) not to be considered in marima.
reg.var	= no. of variable(s) that can only act as regression variable(s) such as (typically) a socalled leading indicator.
no.dep	= sequence of pairs of variables. For example no.dep=c(1, 2, 2, 3) means that variable 2 is not allowed in model for variable 1, and variable 3 is not allowed in model for variable 2.
print	(!0/0) If !0 is used, the generated patterns of the arma model and other informations are printed on the console. If 0 is used, no printout of the arma patterns are given.
ar.fill	= sequence of triplets: c(dependent variable, independent variable, lag). ar.fill=c(2, 3, 12): Insert ar-indicator for model for dependent variable 2 and independent variable 3 at lag 12.
ar.rem	= sequence of triplets c(dependent variable, independent variable, lag). ar.rem=c(2, 3, 12): remove (if present) ar-indicator for model for dependent variable 2 and independent variable 3 at lag 12.
ma.fill	=sequence of triplets: c(dependent variable, independent variable, lag). ma.fill=c(2, 3, 12): Insert ma-indicator for model for dependent variable 2 and independent variable 3 at lag 12.
ma.rem	= sequence of triplets c(dependent variable, independent variable, lag). ma.rem=c(2, 3, 12): remove (if present) ma-indicator for model for dependent variable 2 and independent variable 3 at lag 12.

The various parameters may (in some cases) accomplish the same model requirements. The routine define.model apply these input parameters successively

in the following order: 1) rem.var, 2) reg.var, 3) indep, 4) no.dep, 5) ar.fill, 6) ar.rem, 7) ma.fil, 8) ma.rem

The parameters ar.fill, ar.rem, ma.fill and ma.rem are applied last, and in that order. They overwrite what previously has been defined.

**indep** = no. of variable(s) that are independent of the other variables. indep=c(2, 4) makes variables 2 and 4 independent of all other variables. Variables 2 and 4 may influence other variables.

## Value

ar.pattern = a matrix polynomial with 1's and 0's defining the autoregressive matrix polynomial to be fitted by marima (type=array with dim=c(kvar, kvar, 1+ar\_order) (with leading unity matrix)).

ma.pattern = a matrix polynomial with 1's and 0's defining the moving average matrix polynomial to be fitted by marima (type=array with dim=c(kvar, kvar, 1+ma\_order) (with leading unity matrix)).

## Examples

```
#
# Example 1: 3-variate arma model with ar-lags at 1 and 2, and an
# ma-term at lag 1. And var=3 is a regression variable (X-variable).
#
Model1<-define.model(kvar=3, ar=c(1, 2), ma=c(1), reg.var=3)
short.form(Model1$ar.pattern)
short.form(Model1$ma.pattern, leading=FALSE)
#
# The object Model1 contains the ar- and ma-pattern arrays as defined.
#
# Model1$ar.pattern and Model1$ma.pattern are used as input to
# marima in order to define the model to be estimated.
#
# Example 2: arma model with ar-lags at 1, 2 and 6, and var=3
# regression variable (X-variable).
#
Model2<-define.model(kvar=3, ar=c(1, 2, 6), ma=c(1), reg.var=3)
# Print the ar- and ma-polynomial patterns using
short.form(Model2$ar.pattern, leading=FALSE)
short.form(Model2$ma.pattern, leading=TRUE)
#
# Example 3: arma model with ar-lags at 1, 2 and 6, and reg.var=3
# (X-variable). ma-order=1. Finally (ar.fill=c(2, 3, 4) puts a '1'
# for (dep-var=2, indep-var=3, ar-lag=4).
#
# If further modifications of the ar- or ma-patterns are needed, it
# can be accomplished before calling marima (Model3$ar.pattern and
# Model3$ma.pattern are arrays).
#
Model3<-define.model(kvar=3, ar=c(1, 2, 6), ma=c(1), reg.var=3,
  ar.fill=c(2,3,4))
short.form(Model3$ar.pattern)
short.form(Model3$ma.pattern)
```

```
#
Model4<-define.model(kvar=3, ar=c(1, 2, 6), ma=c(1), reg.var=3,
ar.fill=c(2, 3, 4), indep=c(1))
short.form(Model4$ar.pattern)
short.form(Model4$ma.pattern, leading=FALSE)
```

---

define.sum

*define.sum*


---

## Description

Function to aggregate multivariate time series. Reverse of function 'define.dif'.

## Usage

```
define.sum(series = NULL, difference = NULL, averages = 0)
```

## Arguments

series	= series to be summed up.
difference	= differencing pattern (see define.dif).
averages	of the individual series that (usually) have been subtracted when differencing the time series (if so, the averages are supplied in the output from define.dif(...)).

## Value

sum.series = the summed series.

## Examples

```
set.seed(4711)
y<-round(matrix(100*rnorm(48), nrow=4))
difference=matrix(c(1, 1, 1, 1, 2, 1, 3, 6), nrow=2)
dy<-define.dif(y, difference)$dif.series
averages<-define.dif(y, difference)$averages
sum.y<-define.sum(dy, difference, averages)$series.sum
y
dy
averages
sum.y
```

---

forec.var	<i>forec.var</i>
-----------	------------------

---

**Description**

Function for calculation of variances of nstep forecasts using a marima type model.

**Usage**

```
forec.var(marima, nstep = 1, dif.poly = NULL)
```

**Arguments**

marima	= marima object (cov.u and ar.estimate and ma.estimate are used)
nstep	= length of forecast
dif.poly	= autoregressive representation of differencing polynomial as constructed by the function define.dif(...) when the time series is differenced (if so) before being analysed by marima.

**Value**

= pred.var = variance-covariances for nstep forecasts (an array with dimension (kvar,kvar,nstep)).  
 = rand.shock = corresponding random shock representation of the model used.

---

inverse.form	<i>inverse.form</i>
--------------	---------------------

---

**Description**

Calculation of inverse form for arma model

**Usage**

```
inverse.form(ar.poly, ma.poly, L)
```

**Arguments**

ar.poly	=autoregressive matrix part of model (array(k,k,ar-order)).
ma.poly	=moving average matrix part of model (array(k,k,ma-order)).
L	=order of return polynomial (length=L+1 including leading unity matrix).

**Value**

inverse form for arma model up to order L (array(k,k,L+1)).

## Examples

```
set.seed(4711)
p1 <- check.one(matrix(rnorm(16),nrow=4))
p2 <- check.one(array(rnorm(32),dim=c(4,4,2)))
inverse <- inverse.form(ar.poly=p1,ma.poly=p2,L=6)
short.form(inverse)
```

---

marima

---

*marima*


---

## Description

Estimate multivariate arima and arima-x models. Setting up the proper model for (especially) arima-x estimation can be accomplished using the routine 'define.model' that can assist in setting up the necessary autoregressive and moving average patterns used as input to 'marima'.

A more elaborate description of 'marima' and how it is used can be downloaded from:

<http://www.imm.dtu.dk/~hspl/marima.use.pdf>

## Usage

```
marima(DATA = NULL, ar.pattern = NULL, ma.pattern = NULL, means = 1,
       max.iter = 50, penalty = 0, weight = 0.33, Plot = "none",
       Check = FALSE)
```

## Arguments

DATA	time series matrix, $\text{dim}(\text{DATA}) = c(\text{kvar}, n)$ , where 'kvar' is the dimension of the time series and 'n' is the length of the series. If DATA is organized (n, kvar) (as a data.frame e.g.) it is automatically transposed in marima, and the user need not care about it. Also, and consequently, the output residuals and fitted values matrices are both organised $c(\text{kvar}, n)$ at return from marima. The DATA is checked for completeness. Cases which include 'NA's or 'NaN's are initially left out. A message is given (on the console) and the active cases are given in the output object (...\$used.cases).
ar.pattern	autoregressive pattern for model (see define.model). If ar.pattern is not specified a pure ma-model is estimated.
ma.pattern	moving average pattern for model (see define.model). If ma.pattern is not specified a pure ar-model is estimated. In this case the estimation is carried by regression analysis in a few steps.
means	0/1 indicator vector of length kvar, indicating which variables in the analysis should be means adjusted or not. Default: means=1 and all variables are means adjusted. If means=0 is used, no variables are means adjusted.
max.iter	max. number of iterations in estimation (max.iter=50 is default which, generally, is more than enough).

penalty	parameter used in the R function 'step' for stepwise model reduction. If penalty=2, the conventional AIC criterion is used. If penalty=0, no stepwise reduction of model is performed. Generally $0 \leq \text{penalty} \leq 2$ works well (especially penalty=1). The level of significance of the individual parameter estimates in the final model can be checked by considering the (approximate) 'ar.pvalues' and the 'ma.pvalues' calculated by marima.
weight	weighting factor for smoothing the repeated estimation procedure. Default is weight=0.33 which often works well. If weight>0.33 (e.g. weight=0.66) is specified more damping will result. If a large damping factor is used, the successive estimations are more cautious, and a slower (but safer) convergence (if possible) may result (max.iter may have to be increased to, say, max.iter=75).
Plot	'none' or 'trace' or 'log.det' indicates a plot that shows how the residual covariance matrix (resid.cov) develops with the iterations. If Plot= 'none' no plot is generated. If Plot= 'trace' a plot of the trace of the residual covariance matrix versus iterations is generated. If Plot='log.det' the log(determinant) of the residual covariance matrix (resid.cov) is generated. Default is Plot= 'none'.
Check	(TRUE/FALSE) results (if TRUE) in a printout of some controls of the call to arima. Useful in the first attempt(s) to use marima. Default=FALSE.

## Value

Object of class marima containing:

N = N length of analysed series

kvar = dimension of time series (all random and non-random variables).

ar.estimates = ar-estimates

ma.estimates = ma-estimates

ar.fvalues = ar-fvalues (approximate)

ma.fvalues = ma-fvalues (approximate)

ar.pvalues = ar-p-values (approximate)

ma.pvalues = ma-p-values (approximate)

residuals = estimated residuals (for used.cases)

fitted = estimated/fitted values for all data (including non random variables) (for used.cases)

resid.cov = covariance matrix of (all) residuals (including non random variables) (for used.cases)

data.cov = covariance matrix of (all) input data (for used.cases)

averages = averages of input variables

Constant = estimated model constant =  $(\sum_i(\text{ar}[i, i])) \times \text{averages}$

call.ar.pattern = calling ar.pattern

call.ma.pattern = calling ma.pattern

out.ar.pattern = resulting ar.pattern (after possible model reduction)

out.ma.pattern = resulting ar.pattern (after possible model reduction)

max.iter = max no. of iterations in call

penalty = factor used in AIC model reduction  
 weight = weighting of successive residuals updating (default=0.33)  
 used.cases = cases in input which are analysed  
 trace = trace(random part of resid.cov)  
 log.det = log(det(random part of resid.cov))  
 randoms = which are random variables in problem?

### Source

The code is an R code which is based on the article (below) by Spliid (1983). A repeated (socalled) pseudo regression procedure is used in order to estimate the multivariate arma model.

### References

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Spliid,H.: Estimation of Multivariate Time Series with Regression Variables:  
<http://www.imm.dtu.dk/~hspl/marima.use.pdf>  
[www.itl.nist.gov/div898/handbook/pmc/section4/pmc45.htm](http://www.itl.nist.gov/div898/handbook/pmc/section4/pmc45.htm)

### Examples

```
# Example 1:
library(marima)
# Generate a 4-variate time series (in this example):
#
kvar<-4 ; set.seed(4711)
y4<-matrix(round(100*rnorm(4*1000, mean=2.0)), nrow=kvar)
# If wanted define differencing of variable 4 (lag=1)
# and variable 3 (lag=6), for example:
y4.dif<-define.dif(y4, difference=c(4, 1, 3, 6))
# The differenced series will be in y4.dif$y.dif, the observations
# lost by differencing being excluded.
#
y4.dif.analysis<-y4.dif$y.dif
# Give lags the be included in ar- and ma-parts of model:
#
ar<-c(1, 2, 4)
ma<-c(1)
# Define the multivariate arma model using 'define.model' procedure.
```

```

# Output from 'define.model' will be the patterns of the ar- and ma-
# parts of the model specified.
#
Mod <- define.model(kvar=4, ar=ar, ma=ma, reg.var=3)
arp<-Mod$ar.pattern
map<-Mod$ma.pattern
# Print out model in 'short form':
#
short.form(arp)
short.form(map)
# Now call marima:
Model <- marima(y4.dif.analysis, ar.pattern=arp, ma.pattern=map,
                penalty=0.0)
# The estimated model is in the object 'Model':
#
ar.model<-Model$ar.estimates
ma.model<-Model$ma.estimates
dif.poly<-y4.dif$dif.poly # = difference polynomial in ar-form.
# Multiply the estimated ar-polynomial with difference polynomial
# to compute the aggregated ar-part of the arma model:
#
ar.aggregated <- pol.mul(ar.model, dif.poly, L=12)
# and print everything out in 'short form':
#
short.form(ar.aggregated, leading=FALSE)
short.form(ma.model, leading=FALSE)

```

---

pol.inv

---

*pol.inv*


---

## Description

Calculation of left inverse of matrix polynomial. The leading term is expected to be the (k by k) identity matrix. This is checked and the proper leading unity term is taken into account when the inverse is calculated.

$\phi$  = matrix polynomial coefficients = I,  $\phi_1$ ,  $\phi_2$ , ...,  $\phi(p)$ .

$\dim(\phi) = c(k, k, p+1)$  where k = dimension of coefficient matrices (k by k), and L = order of polynomial (length = 1+L, including the leading unity matrix).

## Usage

```
pol.inv( $\phi$ , L)
```

## Arguments

$\phi$	polynomial to invert
L	order of inverse polynomial



**Value**

left inverse of phi of order L (L+1 terms including leading unity matrix)

**Examples**

```
set.seed(4711)
p2<-check.one(array(rnorm(32),dim=c(4,4,2)))
pi2<-pol.inv(p2,L=12)
short.form(pi2)
```

---

<code>pol.mul</code>	<i>pol.mul</i>
----------------------	----------------

---

**Description**

Calculation of product of two matrix polynomials (arrays).  
If one or both leading unity matrices (of eta and theta) are missing, they are (it is) generated (and taken into account).

**Usage**

```
pol.mul(eta, theta, L)
```

**Arguments**

- eta                    first matrix polynomial
- theta                second matrix oynomial
- L                    order of output polynomial (length = L+1)

**Value**

matrix polynomial product af eta and theta

**Examples**

```
set.seed(4711)
p1<-check.one(matrix(rnorm(16),nrow=4))
p2<-check.one(array(rnorm(32),dim=c(4,4,2)))
p12<-pol.mul(p1,p2,L=(2+3))
short.form(p12)
```

---

pol.order	<i>pol.order</i>
-----------	------------------

---

**Description**

Function to evaluate (significant) order of matrix polynomial.

**Usage**

```
pol.order(polyn = NULL, digits = 12)
```

**Arguments**

polyn	the polynomial the order of which is determined.
digits	number of significant digits to be considered (values smaller than $10^{-(\text{digits})}$ are taken to be 0 (zero)).

**Value**

pol.order order of polynomial polyn. (exclusive the leading unity matrix if present. pol.order=0 corresponds to the k by k unity matrix)

**Examples**

```
pol      <- array(1e-8*rnorm(96),dim=c(4,4,6))
pol[, , 1:3] <- array(rnorm(48),dim=c(4,4,3))
pol.order(polyn=pol,digits=12)
pol.order(polyn=pol,digits=4)
```

---

print.marima	<i>print.marima</i>
--------------	---------------------

---

**Description**

Print some (most relevant) content of a marima object.

**Usage**

```
## S3 method for class 'marima'
print(x, estimates = TRUE, pvalues = FALSE,
      pattern = TRUE, fvalues = TRUE, ...)
```

**Arguments**

x	= a marima object with results of marima analysis.
estimates	= TRUE/FALSE: printout of parameter estimates.
pvalues	= TRUE/FALSE: printout of (approximate) p-values for parameter estimates.
pattern	= TRUE/FALSE: printout of model definition pattern(s).
fvalues	= TRUE/FALSE: printout of parameter (approximate) F-values.
...	Not used.

---

rand.shock

*rand.shock*


---

**Description**

Calculation of random shock form for arma model

**Usage**

```
rand.shock(ar.poly, ma.poly, L)
```

**Arguments**

ar.poly	autoregressive matrix part of model
ma.poly	moving average matrix part of model
L	order of return polynomial (length=L+1 including leading unity matrix)

**Value**

random shock form of arma model up to order L (array(k,k,L+1))

**Examples**

```
set.seed(4711)
p1 <- check.one(matrix(rnorm(16),nrow=4))
p2 <- check.one(array(rnorm(32),dim=c(4,4,2)))
randshock <- rand.shock(ar.poly=p1,ma.poly=p2,L=6)
short.form(randshock)
```

---

season.lagging	<i>season.lagging</i>
----------------	-----------------------

---

## Description

Generate new time series with (seasonally) lagged variables from lagging pattern.

## Usage

```
season.lagging(y, lagging = NULL)
```

## Arguments

**y** = data series

**lagging** = lagging array array describing what to be added to y: c(1, 3, 6) adds a new y3, using y1 lagged 6 time steps. lagging<-matrix(c(1, 3, 6-1, 2, 4, 12-1), nrow=3) adds two new variables (y3 and y4) using y1 lagged 6-1 time steps and y2 lagged 12-1 time steps.

## Value

y.lagged = the part of the new series (including new lagged variables) that can be entered into marima

y.future = the part of the new series (including new lagged variables) that does not include future observation

y.lost = previous values of the time series that is incomplete with respect to the new variables generated by lagging

cbind(y.lost, y.lagged.y, y.future) is the complete series after creation and addition of the lagged variables.

## Examples

```
set.seed(4711)
# generate bivariate time series
y<-round(matrix(10*rnorm(36), nrow=2))
y
# define new lagged variables (y3 and y4) with seasonalities 6 and 12
lagging <-c(1, 3, (6-1), 2, 4, (12-1)) #
season.lagging(y, lagging)
```

---

short.form

*short.form*


---

## Description

Function to condensate (and/or) print out matrix polynomial leaving out empty lag matrices and, if specified, the leading (unity) matrix.

## Usage

```
short.form(poly = NULL, name = "Lag=", leading = TRUE, tail = FALSE,
  digits = 6)
```

## Arguments

poly	matrix polynomial (0-1 array as constructed by define.model, for example, or array of reals as estimated by marima).
name	character string used as header in output (default='lag').
leading	TRUE/FALSE. If leading=FALSE the leading (unity matrix) is to be left out/suppressed.
tail	TRUE/FALSE. If TRUE and the ar/ma-model only consists of coefficient matrix(s) where all coefficients (except the leading unity matrix) are all zero a first order coefficient matrix (being zero) is retained (in order to avoid a model containing only the leading unity matrix). If tail=TRUE and the coefficients in the first coefficient matrix (after the leading unity matrix) are all zero, the leading unity matrix is always retained.
digits	the number of digits retained by short.form (default=6).

## Examples

```
Model<-define.model(kvar=4, ar=c(1, 2, 4), ma=c(1), reg.var=4)
short.form(Model$ar.pattern)
short.form(Model$ma.pattern)
short.form(Model$ar.pattern, leading=FALSE)
short.form(Model$ar.pattern, leading=FALSE)
#
M<-define.model(kvar=4, ma=c(1))
short.form(M$ar.pattern)
short.form(M$ar.pattern, tail=TRUE)
short.form(M$ar.pattern, leading=FALSE, tail=TRUE)
```

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